

**● PRINTER RUSH ●**  
**(PTO ASSISTANCE)**

Application : <u>09/744,167</u>	Examiner : <u>Borin</u>	GAU : <u>1631</u>
From: <u>J. Robbins</u>	Location: <u>(IDC) FMF FDC</u>	Date: <u>1-11-06</u>
Tracking #: <u>06041103</u>		Week Date: <u>11-15-04</u>

DOC CODE	DOC DATE	MISCELLANEOUS
<input type="checkbox"/> 1449	_____	<input type="checkbox"/> Continuing Data
<input type="checkbox"/> IDS	_____	<input type="checkbox"/> Foreign Priority
<input type="checkbox"/> CLM	_____	<input type="checkbox"/> Document Legibility
<input type="checkbox"/> IIFW	_____	<input type="checkbox"/> Fees
<input type="checkbox"/> SRFW	_____	<input checked="" type="checkbox"/> Other <u>Let. 11-3-2005</u>
<input type="checkbox"/> DRW	_____	
<input type="checkbox"/> OATH	_____	
<input type="checkbox"/> 312	_____	
<input type="checkbox"/> SPEC	_____	

**[RUSH] MESSAGE:** Page 62, Tables 9 and 10 which were provided per previous response to XRUSH. (See Let. 11-3-2005) are still illegible for publication

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**[XRUSH] RESPONSE:** \_\_\_\_\_

Done

clearest copy provided

**INITIALS:** [Signature]

NOTE: This form will be included as part of the official USPTO record, with the Response document coded as XRUSH.  
REV 10/04

Table 9

hSARA	MFNDEHVAEDQLIKRNYSDDCSAVEGERKCGMLACLPDEKNVLVVAVHHNCDKRTQNDLQDCNNYNSQSLEND	77
XSARA	MPKVVGGDTTMAEDSLFNNIGPSEIVNPSTVSSG---SLQALDDQS---VNIHNEKSVILLADGFSPSSP---KSIIN	70
hSARA	AFSLDNEHQTDQFSSINESIEKDNINSEKNDPLNRPKTEGRSVNHCPTSSSLASVCSFSQLDDGSIERDPSMS	157
XSARA	FDCTHMDN-----EMLHSONMSVDNDN---ETVITISVLPTIIQDTSNVSDPAINKP---GPKEPHR	127
hSARA	ATTSITVDSVISSCGTDGFAVKKQEN-YLPDELTGKISSPRTDLGSPNSFSHASEGILMKKEPAEESTTEESIRSSLP	236
XSARA	ALKETTSVILPEIKPYSTCAALSFERANKVPSYQLN-----NTDILS---VSPVVEACSEQQQKHTSSLHEEKLPFGVS	198
hSARA	LLLKPDPNPGSGRNDCCERCSDCLVPNEVRADENEGYEHEETLCTTEFLNNTTEFSESQDMTKMLTKLNEFNDSQVNEE	316
XSARA	ATES-----FAATAETVLONEALSAEFTDIUVKNFSDSCVINGDLTKSCGLS---QES	250
hSARA	KEKFLQISPEEDTNGDSGGQCVGLADAGLDDGTCTISESECDPSTVIDTPAANVLSNGCDSYGMQDPCGVSEVPKTLPSK	396
XSARA	NEKFCASKEPF-----CG-----VDANVLLNACVAYKEADLPPEENGQINAPMELLYNGCDSYGMKNPFAVAQNPKNLPSK	319
hSARA	EDSVTEEKEIEESKSEYSNIEYEQFGNEATESSGLLNSTCDLHKKNYLMHFCQVPSVLGQSSPKVVASLPSISVFPFG	475
XSARA	EDSVTEEKEIEESKSEYVTGVYEQDFEDVTERGGLLLNAKADQMKNN-LHSLOQVPSMAGQTSPPKGGKIVQSLSVPYG	398
hSARA	GARPQPSNLKLQIPKPLSDHLND-FFANSGNNTTKKNDILGKAKLGENSATNVCSPLIGNISNVDPTNGEHLESYEAE	553
XSARA	GARTKQPTHKLHHPKPLSEMLQSLDIPNACSSKYKNDHMLNKSNOGDMLISESRLREDSAVRSPVTDANGDFPGEYRGP	478
hSARA	ISTRPLALAPDSPDNOLAGQFGISARKPFTTLGEVAPVWVPDSQAPNCKMCEARFTFTKRHHCRACGKVFCASCCSL	633
XSARA	GS-ICLAVSPDSPDNOLAGQFGVPIISKPFTTLGEVAPVWVPDSQAPNCKMCEARFTFTKRHHCRACGKVFCACCCSL	556
hSARA	KCKILYMDRKEARVCVICHSVLMNAQAWENHMSASSOSPNNPAEYCSPTIPPLQQAQASGALSSPPPTVNVPGVLKHP	713
XSARA	KCKILOYMDKKEARVCVICHSVLMNAQAWENMLSASVOSPNNPAEYCSPTIPPMQQAQASGALSSPPPTVNVPGVLKHP	636
hSARA	GAEVACREQRRVWFADGILPNGEVADAAKITNGCTSSAGTLAVSHDPVKPVITISPLPAETDITLFSGSITQVGSVPVGEA	793
XSARA	GTESSCKEORRVWFADGILPNGETADSD--NNNVITVAGTLTVSHNNNSTSSCS-----ENTSGFCGSITQVG---SA	705
hSARA	MNLIPEDGLPILISTGVKGDYAVEEKPSQISVMHQLEDGGPDPLVFLNANLLSHVKIVNYVNRKWCFTTKGHHAVGQ	873
XSARA	MNLIPEDGLPILISTGVKGDYAVEERPQMSVMHQLEEGPDPLVFLNANLLAHVKIVNYVNRKWCFTTKGHHAVGQ	785
hSARA	SEIVILLQCLPDEKCLPKDIEHMFVQLYRDALAGNVVNLGHSFESQSFLGSKHEGGFLYVITSTYQSLQDLVLPTPEYLF	953
XSARA	AEIVILLQCLPDEKCLPRDLESHFVELYQEAIGAGNVVNLGHSFESQSFLGSKDHGGFLYVAFTYQSLQDLVLPAPEYLF	865
hSARA	GILIQKWETPWAKVFFIRLNLRLGAEYRLYPCLFSVRFRKPLPGETGHTIMNLLADFRNYQYTLPPVQGLVVDMEVRKT	1033
XSARA	GILIQKWETPWAKVFFIRLNLRLGAEYRLYPCLFSVRYRKPLPGETGHTIINVLADFRNYQYTLPPVQGLVVDMEVRKT	945
hSARA	SIKIPSNRYNEMHKAMNKSNEHVLAAGACFNEKADSHLVCVQNDGNYQTQAISIHQPRKVTGASFFVFSGALKSSSGY	1113
XSARA	SIKIPSNRYNEMHKAMNKSNEHVLAAGACFNOADSHLVCVQNDGNYQTQAISIHQPRKVTGASFFVFSGALKSSSGY	1025
hSARA	LAKSSIVEDGVHVQITAEIMDSLRQALREMKDFTITCGKADAEEQEHVHIQWVDDKMYSGKGVSPIDGKSMETITNVK	1193
XSARA	LAKSSIVEDGVHVQITAEIMDALRQSLREMKDFTITCGKADAEEQEHVHVQWVEDDKMYSGKGVSPIDGKSMESVTSVK	1105
hSARA	IFHGSEYKANKKVIKRMTEVFFLENDQHNCLSDPADHSRLTEHVAKAFCLALCPHLKLLKEDGMTRLGLRVTLDSQVGY	1273
XSARA	IFHGSEYKASGKIIRMTDEVFFLDNEEQCSGLSDPADHSRLTENVAKAFCLALCPHLKLLKEDGMTRLGLRVSLDSQVGY	1185
hSARA	QAGSNGQFLPSQYINDLDSALVPVTHGGACQLSEGPVVMELIFYILENTV- 1323	
XSARA	QAGSNGQILPARYINDLDSALVPVTHGGTCQLSEGPVVMELIFYILENTS- 1235	

Table 10

hSARA	587	GEVAPVWVPDSQAPNCKMCEARFTFTKRHHHCRACGKVFCASCCSLKCKLYMDRKEARVCVICHSVI	655
XSARA	510	LGEVAPVWVPDSQAPNCKMCEARFTFTKRHHHCRACGKVFCACCSLKCKLYMDRKEARVCVICHSVI	578
KIAA0305	737	LGCKDPVWVPDSQAPNCKMCEARFTFTKRHHHCRACGKVFCACCSLKCKLYMDRKEARVCVICHSVI	800
FGD1	720	LGRAPITRIPEKEVITCMHCQEPFNSITKRHHHCRACGKVFCACCSLKCKLYMDRKEARVCVICHSVI	485
Hrs	153	AAERAPVWD- -AEECHRCRVQF- GVVTRKHHCRACGQIFCGKCSSKYSTIPKFGIEKEVRVCEPCYEOL	219
Hrs-2	153	AAERAPVWD- -AEECHRCRVQF- GVVTRKHHCRACGQIFCGKCSSKYSTIPKFGIEKEVRVCEPCYEOL	219
EEA-1	1341	TOALNRKMAEDNEVONCMACGKEF-SVTVRRHHCRACGQIFCAECSAKNALTPSSKIL- PVAVCDACFNDL	1408
CONSENSUS - - - - P-W - - - - - C-C - - - F - - - - R <sup>R</sup> KHHCRACG-YFC <sup>A</sup> CS <sup>S</sup> - - - - - RVC - - C - - L			